7,1400

SEQUENCE LISTING

<110>	Gao, Zeren Hart, Charles E. Piddington, Christopher S. Sheppard, Paul O. Shoemaker, Kimberly E. Gilbertson, Debra G. West, James W.
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	tct gcc ctg gcc ggc cag aga cag ggg act cag gcg gaa Ser Ala Leu Ala Gly Gln Arg Gln Gly Thr Gln Ala Glu 15 20
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		cag Gln 45									318
		cca Pro									366
		tta Leu					_				414
		aga Arg		-	_		-	-	_		462
		gta Val									510
		ggt Gly 125									558
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		tgc Cys				-	-				654
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	tcc Ser															894
	agc Ser															942
	aga Arg 265		-							-		-	-		•	990
	ggt Gly													_		1038
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	ctg Leu								_		_	_		-		1182
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gcti agaa ctta ttaa ggta aact ttti	caag attag caatg catg aatgt	gga ogga g gga g gta og ca g aag o	cttt gttgt ggaaa gtat gtaca ctcca tatta	cato gcaa agaaa tcca aggaa atgto cacat	et to accompany to the	agga jctct aaat igctg aact jggco jtaaa	attta ttttg tgttg gggtt gggg taaa	a cag g aga g tat c ctg c aag a ato g aac	gtgca aggag taat gtatt gtgag cgtat	attc ggcc atag atca gcac aaa atat	tgaa taaa atca gtto ctga atct gtao	agagagagagagagagagagagagagagagagagagag	gga g cag g gct a cga t cgt t ctt t	gacat gagaa agttt tacgg tgcct	ctgttt ccaaac aaaggt ccagag gcttag ctgctt cttttt	1291 1351 1411 1471 1531 1591 1651 1711 1760



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Tyr His Glu Val Leu Gln Leu Arg Pro Lys Thr Gly Val Arg Gly Leu 305 310 315 320 His Lys Ser Leu Thr Asp Val Ala Leu Glu His His Glu Glu Cys Asp

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gtncargayc cncarcayga rmgnathath acngtnwsna cnaayggnws nathcaywsn	180
conmunity concayaonta yoonmgnaay acnginying iniggmgnyi ngingongin	240
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carytnytng gnaargcntt ygtnttyggn mgnaarwsnm gngtngtnga yytnaayytn	720
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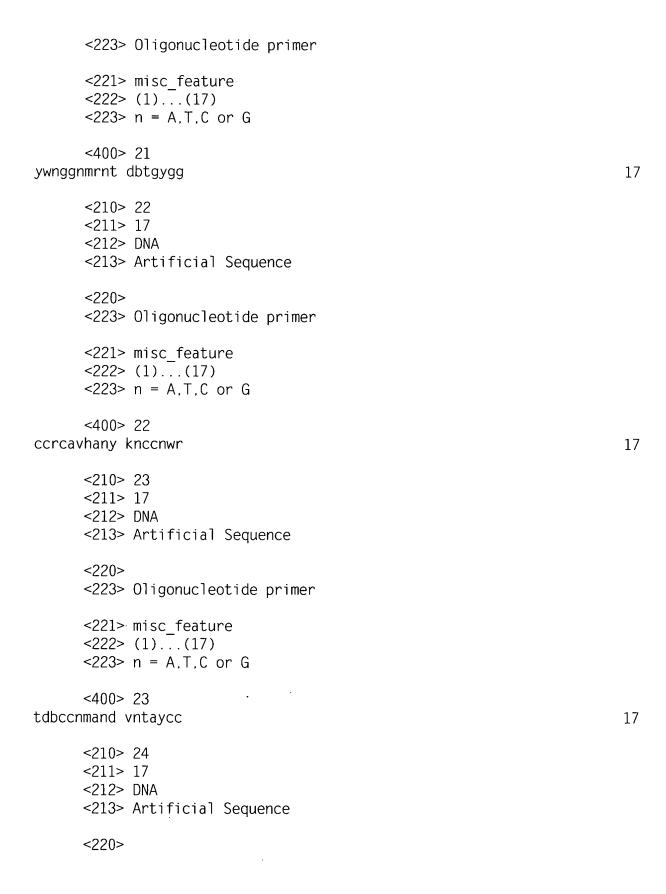
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Glu Asp Ile Ser Glu Thr Ser Thr Ile Ile Arg Gly Arg Trp Cys Gly

125

120



130

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												ttc Phe	-	717
												tca Ser		765
					_					-		tat Tyr 195		813
	_	_	-			-			_	-	•	gac Asp		861
			_			-	_	_		-		ttc Phe		909
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												gtc Val		1101
												ggc Gly		1149



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agg ggt aga gct aag acc atg gct cta gtt gac atc cag ttg gat cac Arg Gly Arg Ala Lys Thr Met Ala Leu Val Asp Ile Gln Leu Asp His 345 350 355	1293
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Pro Gly Phe Lys Ile Tyr Tyr Ser Leu Leu Glu Asp Phe Gln Pro Ala
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Leu Asp Thr Pro Arg Tyr Arg Gly Arg Ser Tyr His Asp Arg Lys Ser
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Thr Pro Arg Asn Tyr Ser Val Asn Ile Arg Glu Glu Leu Lys Leu Ala
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agc atc cac agc ccg a Ser Ile His Ser Pro L 60	=	s Thr Tyr Pr	_	
ctg gtg tgg aga tta g Leu Val Trp Arg Leu V 75				
aca ttt gat gag aga t Thr Phe Asp Glu Arg P 90			lu Asp Asp Il	



_	tat Tyr	-	_	_	_				_	-		-	-		1408
	cgc Arg														1456
	aat Asn				-		-		-						1504
	ccc Pro														1552
	acc Thr 170														1600
_	ctc Leu		_			-		_		_	-		_		1648
	tac Tyr														1696
	cca Pro														1744
	aaa Lys														1792
_	tgc Cys 250										_	-		-	1840
	aca Thr	_						-		Leu	_	_	-	-	1888



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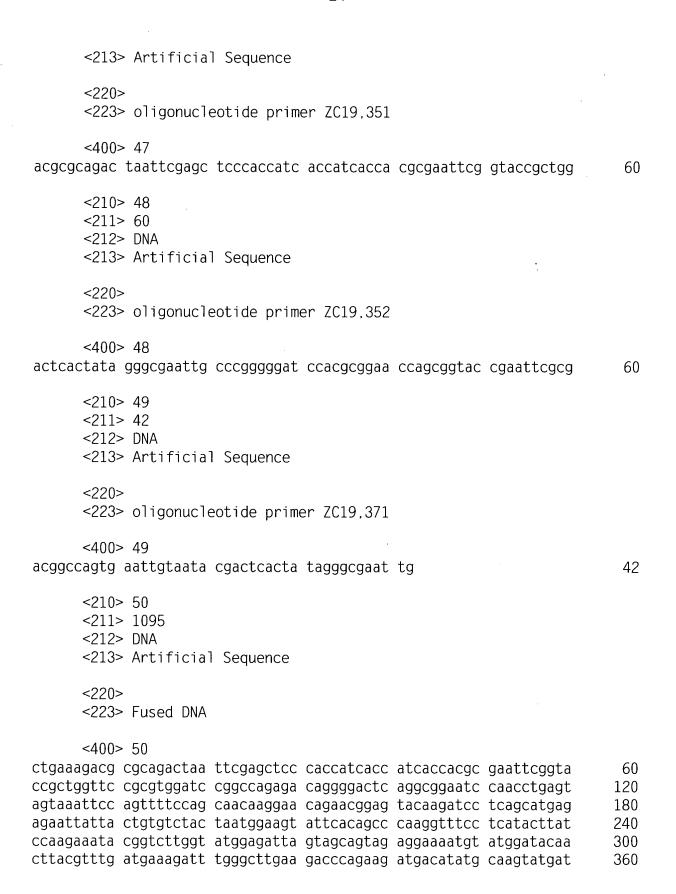
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295

164

Tyr 305	His	Glu	Val	Leu	Gln 310	Leu	Arg	Pro	Lys	Thr 315	Gly	Val	Lys	Gly	Leu 320		
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(66





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